

GenCore version 5.1.4_P5_4578
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OM protein - protein search, using SW model

Run on: April 1, 2003, 08:47:16 ; Search time 18 Seconds
(without alignments)
1525.005 Million cell updates/sec

Title: US-09-768-781-3
Perfect score: 2316
Sequence: 1 MDRVYETIPEEPNVDPVSSLE.....RTRVENSEPPFTEARRQSVV 449

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 248812 seqs, 61136040 residues

Total number of hits satisfying chosen parameters: 248812

Minimum DB seq length: 0
Maximum DB seq length: 2000000000
Post-processing: Minimum Match 0%
Maximum Match 100%

Listing First 45 summaries

Database : Published Applications AA:*

1: /cgnd2_6/_ptodata/1/pubpaas/us08_NEW_PUB.pep:*

2: /cgnd2_6/_ptodata/1/pubpaas/us01_NEW_PUB.pep:*

3: /cgnd2_6/_ptodata/1/pubpaas/us06_NEW_PUB.pep:*

4: /cgnd2_6/_ptodata/1/pubpaas/us06_NEW_PUBCOMB.pep:*

5: /cgnd2_6/_ptodata/1/pubpaas/us07_NEW_PUB.pep:*

6: /cgnd2_6/_ptodata/1/pubpaas/us07_PUBCOMB.pep:*

7: /cgnd2_6/_ptodata/1/pubpaas/pctus_pubcomb.pep:*

8: /cgnd2_6/_ptodata/1/pubpaas/us08_PUBCOMB.pep:*

9: /cgnd2_6/_ptodata/1/pubpaas/us09_NEW_PUB.pep:*

10: /cgnd2_6/_ptodata/1/pubpaas/us09_NEW_PUBCOMB.pep:*

11: /cgnd2_6/_ptodata/1/pubpaas/us10_NEW_PUB.pep:*

12: /cgnd2_6/_ptodata/1/pubpaas/us10_PUBCOMB.pep:*

13: /cgnd2_6/_ptodata/1/pubpaas/us60_NEW_PUB.pep:*

14: /cgnd2_6/_ptodata/1/pubpaas/us60_PUBCOMB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

| Result No. | Score | Query Match | Length | DB ID | Description |
|------------|-------|-------------|--------|-----------------------|-------------------|
| 1 | 2316 | 100.0 | 449 | 10 US-09-768-781-3 | Sequence 3, Appli |
| 2 | 2301 | 99.4 | 449 | 10 US-09-768-781-7 | Sequence 7, Appli |
| 3 | 2301 | 99.4 | 462 | 10 US-09-768-781-4 | Sequence 4, Appli |
| 4 | 939.5 | 40.6 | 405 | 10 US-09-768-781-6 | Sequence 6, Appli |
| 5 | 607 | 24.7 | 216 | 9 US-09-864-161-3387 | Sequence 33387, A |
| 6 | 572 | 24.7 | 125 | 9 US-10-092-154-595 | Sequence 595, App |
| 7 | 572 | 24.7 | 125 | 10 US-09-744-847-595 | Sequence 595, App |
| 8 | 354 | 15.3 | 128 | 10 US-10-175-73-524 | Sequence 37894, A |
| 9 | 174.9 | 7.5 | 86 | 10 US-09-864-761-3764 | Sequence 35764, A |
| 10 | 113 | 4.9 | 686 | 9 US-10-174-590-524 | Sequence 524, App |
| 11 | 113 | 4.9 | 686 | 9 US-10-176-578-524 | Sequence 524, App |
| 12 | 113 | 4.9 | 686 | 9 US-10-175-37-524 | Sequence 524, App |
| 13 | 113 | 4.9 | 686 | 9 US-10-173-706-524 | Sequence 524, App |
| 14 | 113 | 4.9 | 686 | 9 US-10-175-73-524 | Sequence 524, App |
| 15 | 113 | 4.9 | 686 | 9 US-10-175-752-524 | Sequence 524, App |
| 16 | 113 | 4.9 | 686 | 9 US-10-176-482-524 | Sequence 524, App |
| 17 | 113 | 4.9 | 686 | 9 US-10-176-577-524 | Sequence 524, App |
| 18 | 113 | 4.9 | 686 | 9 US-10-176-913-524 | Sequence 524, App |
| 19 | 113 | 4.9 | 686 | 9 US-10-180-552-524 | Sequence 524, App |

ALIGNMENTS

RESULT 1
US-09-768-781-3
; Application US/09768781
; Patent No. US20020142372A1
; GENERAL INFORMATION:
; APPLICANT: MERKULOV, Gennady V. et al
; TITLE OF INVENTION: ISOLATED HUMAN TRANSPORTER PROTEINS,
; NUCLEAR ACID MOLECULES ENCODING HUMAN TRANSPORTER PROTEINS,
; TITLE OF INVENTION: AND USES THEREOF
; FILE REFERENCE: C1001057-CLP
; CURRENT APPLICATION NUMBER: US/09/768-781
; CURRENT FILING DATE: 2001-01-25
; SOFTWARE: RastSeq for Windows Version 4.0
; SEQ ID NO: 7
; SEQ ID NO: 3
; LENGTH: 449
; TYPE: PRT
; ORGANISM: Human
US-09-768-781-3

| Query Match | Best Local Similarity | Score | DB 10; | DB 11; | DB 12; | DB 13; | DB 14; | DB 15; | DB 16; | DB 17; | DB 18; | DB 19; | DB 20; |
|-------------|-------------------------|----------------------------|---------------------|---------------------|---------------------|---------------------|---------------------|---------------------|---------------------|---------------------|---------------------|---------------------|---------------------|
| 1 | 100.0%; Matches 449; | 100.0%; Conservative 0; | Score 2316; |
| 2 | 100.0%; Matches 449; | 100.0%; Conservative 0; | Pred. No. 3.2e-213; |
| 3 | 100.0%; Matches 449; | 100.0%; Conservative 0; | Mismatches 0; | Mismatches 0; | Mismatches 0; | Mismatches 0; | Mismatches 0; | Mismatches 0; | Mismatches 0; | Mismatches 0; | Mismatches 0; | Mismatches 0; | Mismatches 0; |

SUMMARIES

| Result No. | Score | Query Match | Length | DB ID | Description |
|------------|-------|-------------|--------|-----------------------|-------------------|
| 1 | 2316 | 100.0 | 449 | 10 US-09-768-781-3 | Sequence 3, Appli |
| 2 | 2301 | 99.4 | 449 | 10 US-09-768-781-7 | Sequence 7, Appli |
| 3 | 2301 | 99.4 | 462 | 10 US-09-768-781-4 | Sequence 4, Appli |
| 4 | 939.5 | 40.6 | 405 | 10 US-09-768-781-6 | Sequence 6, Appli |
| 5 | 607 | 24.7 | 216 | 9 US-09-864-161-3387 | Sequence 33387, A |
| 6 | 572 | 24.7 | 125 | 9 US-10-092-154-595 | Sequence 595, App |
| 7 | 572 | 24.7 | 125 | 10 US-09-744-847-595 | Sequence 595, App |
| 8 | 354 | 15.3 | 128 | 10 US-10-175-73-524 | Sequence 37894, A |
| 9 | 174.9 | 7.5 | 86 | 10 US-09-864-761-3764 | Sequence 35764, A |
| 10 | 113 | 4.9 | 686 | 9 US-10-174-590-524 | Sequence 524, App |
| 11 | 113 | 4.9 | 686 | 9 US-10-176-578-524 | Sequence 524, App |
| 12 | 113 | 4.9 | 686 | 9 US-10-175-37-524 | Sequence 524, App |
| 13 | 113 | 4.9 | 686 | 9 US-10-173-706-524 | Sequence 524, App |
| 14 | 113 | 4.9 | 686 | 9 US-10-175-73-524 | Sequence 524, App |
| 15 | 113 | 4.9 | 686 | 9 US-10-175-752-524 | Sequence 524, App |
| 16 | 113 | 4.9 | 686 | 9 US-10-176-482-524 | Sequence 524, App |
| 17 | 113 | 4.9 | 686 | 9 US-10-176-577-524 | Sequence 524, App |
| 18 | 113 | 4.9 | 686 | 9 US-10-176-913-524 | Sequence 524, App |
| 19 | 113 | 4.9 | 686 | 9 US-10-180-552-524 | Sequence 524, App |

Db 241 LCITIWRWLBITSRLLVLFVLSATIKLKAVPFLVINFLLILFEPWIKFWSGAQMPNNIE 300
 Qy 301 KNPSRGVTLYVLISVTIYAGINSQCSWASQLRQLADRLVDKGONGWGLNLYSVRLVEN 360
 Db 301 KNPSRGVTLYVLISVTIYAGINSQCSWASQLRQLADRLVDKGONGWGLNLYSVRLVEN 360
 Qy 361 VIMVLVKEFPGVKLNYCHSLLIAQLIAYLISIDFMLLFFQYLHPLRSLFTNVVDYL 420
 Db 361 VIMVLVKEFPGVKLNYCHSLLIAQLIAYLISIDFMLLFFQYLHPLRSLFTNVVDYL 420
 ; TITLE OF INVENTION: ISOLATED HUMAN TRANSPORTER PROTEINS,
 ; TITLE OF INVENTION: NUCLEAR ACID MOLECULES ENCODING HUMAN TRANSPORTER PROTEINS,
 ; TITLE OF INVENTION: AND USES THEREOF
 ; FILE REFERENCE: CL001057-CLP
 ; CURRENT APPLICATION NUMBER: US/09/768,781
 ; CURRENT FILING DATE: 2001-01-25
 ; NUMBER OF SEQ ID NOS: 7
 ; SOFTWARE: FastSEQ for Windows Version 4.0
 ; SEQ ID NO: 4
 ; LENGTH: 462
 ; TYPE: PRT
 ; ORGANISM: Human

RESULT 2

US-09-768-781-7
 ; Sequence 7, Application US/09768781
 ; Patent No. US20020142376A1
 ; GENERAL INFORMATION:
 ; APPLICANT: MERKULOV, Gennady V. et al
 ; TITLE OF INVENTION: ISOLATED HUMAN TRANSPORTER PROTEINS,
 ; TITLE OF INVENTION: NUCLEAR ACID MOLECULES ENCODING HUMAN TRANSPORTER PROTEINS,
 ; TITLE OF INVENTION: AND USES THEREOF
 ; FILE REFERENCE: CL001057-CLP
 ; CURRENT APPLICATION NUMBER: US/09/768,781
 ; CURRENT FILING DATE: 2001-01-25
 ; NUMBER OF SEQ ID NOS: 7
 ; SOFTWARE: FastSEQ for Windows Version 4.0
 ; SEQ ID NO: 7
 ; LENGTH: 449
 ; TYPE: PRT
 ; ORGANISM: Mus Musculus

US-09-768-781-7

Query Match 99.4%; Score 2301; DB 10; Length 462;
 Best Local Similarity 99.6%; Pred. No. 8.9e-12;
 Matches 447; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 MDRYTYEIPSEPNVDPVSSLEEDVIRGANPRPTFFPSILSTFLYGEASALYMYRIYRK 60
 Db 14 MDRYTYEIPPEPNVDPVSSLEEDVIRGANPRPTFFPSILSTFLYGEASALYMYRIYRK 60
 ;
 Qy 61 NSETYRMYTFSFPFMSSIMVQLTLIIVYRDLAKDKPLSFMLHLLGPVIRCLEAMIKY 120
 Db 74 NSETYWMYTFSFPFMSSIMVQLTLIIVYRDLAKDKPLSFMLHLLGPVIRCLEAMIKY 133
 ;
 Qy 121 LTWKKEKEOEPYPSLTREKMLIDGEVLTWEYGHSLTMLHMRNAYKMSQIOAFLGS 180
 Db 134 LTWKKEKEOEPYPSLTREKMLIDGEVLTWEYGHSLTMLHMRNAYKMSQIOAFLGS 193
 ;
 Qy 181 VPQTYQLVSLISAEPVPLGVVLMVFSLVSVTIGATLCNLMLAQIKYDDYKIRGPLEV 240
 Db 194 VPQTYQLVSLISAEPVPLGVVLMVFSLVSVTIGATLCNLMLAQIKYDDYKIRGPLEV 253
 ;
 Qy 241 LCITIWRWLBITSRLLVLFVLSATIKLKAVPFLVINFLLILFEPWIKFWSGAQMPNNIE 300
 Db 254 LCITIWRWLBITSRLLVLFVLSATIKLKAVPFLVINFLLILFEPWIKFWSGAQMPNNIE 313
 ;
 Qy 301 KNSFRVGTLVVLISVTIYAGINFSCWASQLRQLADRLVDKGONGWGLHYSVRLVEN 360
 Db 314 KNSFRVGTLVVLISVTIYAGINFSCWASQLRQLADRLVDKGONGWGLHYSVRLVEN 373
 ;
 Qy 361 VIMVLVKEFPGVKLNYCHSLLIAQLIAYLISIDFMLLFFQYLHPLRSLFTNVVDYL 420.
 Db 374 VIMVLVKEFPGVKLNYCHSLLIAQLIAYLISIDFMLLFFQYLHPLRSLFTNVVDYL 433
 ;
 Qy 421 HCVCCHOHPRTVENSEPPFETEARQSVV 449
 Db 434 HCVCCHOHPRTVENSEPPFETEARQSVV 462
 ;
 Qy 462 LCITIWRWLBITSRLLVLFVLSATIKLKAVPFLVINFLLILFEPWIKFWSGAQMPNNIE 300
 Db 474 LCITIWRWLBITSRLLVLFVLSATIKLKAVPFLVINFLLILFEPWIKFWSGAQMPNNIE 300
 ;
 Qy 501 KNPSRGVTLYVLISVTIYAGINSQCSWASQLRQLADRLVDKGONGWGLNLYSVRLVEN 360
 Db 501 KNPSRGVTLYVLISVTIYAGINSQCSWASQLRQLADRLVDKGONGWGLNLYSVRLVEN 360
 ;
 Qy 561 VIMVLVKEFPGVKLNYCHSLLIAQLIAYLISIDFMLLFFQYLHPLRSLFTNVVDYL 420
 Db 561 VIMVLVKEFPGVKLNYCHSLLIAQLIAYLISIDFMLLFFQYLHPLRSLFTNVVDYL 420
 ;
 Qy 621 HCVCCHOHPRTVENSEPPFETEARQSVV 449
 Db 621 HCVCCHOHPRTVENSEPPFETEARQSVV 449
 ;
 RESULT 3

RESULT 3

ORGANISM: Mus musculus
US-09-768-781-6

Query Match 40.6%; Score 939.5; DB 10; Length 405;
Best Local Similarity 43.8%; Pred. No. 9.6e-82;
Matches 180; Conservative 85; Mismatches 135; Indels 11; Gaps 4;

Qy 33 PPFSLFSTFLYCGEANSALYMRVYRKNSETYRMTYTFSEPMESSIMVQLTLYFVHDL 92
Db 1 FPASTASVFLVFAETAALYLSSITRSAGDRMWQVLTFLMPCLAVQFTLFLVHDL 60

Qy 93 ARDKPLSLFLMHLILGEVIRCLEAMIXYLTMKKEOBEPYVSLTRKX-MLJDGEELIE 151
Db 61 SRDRPLALLMHLQLGLPLYRCBVFCLYC---QSDQNEBEPYVSITKRCOMPDKGSLREVE 117

Qy 152 WEGHSIERTLAMHNAYKRMOSIQAPLGSPVOLTYQIYVLSIASEVPGLGRVYLMVFSLVS 211
Db 118 KEVGQAEGKLTHRSAFSRASTVQAPLGSAPOLTQLYITVLEQNITTGRCFTMTLSLLS 177

Qy 212 VTYGATLCNMLAIQIKYDDYKIRLGPLBLVKPLVLTQWTRLEITSRLLVLFSATLKURAVP 271
Db 178 IVYGALRCNLIAIKYDYEYVKVPLAYCFLWVCFIWRVTLVPLSVLKWWVA 237

Qy 272 FLVNPFLILIFPEPWKFWRSGAQMPNNIEKNFSRGVTLVVLISVTIYAGINFSCMSALQ 331
Db 238 VILVNPFSFLPYWVWCSSGPPEPIEKALSRVGTTIVLCFLTLYAGINMPCMSAVQ 297

Qy 332 IRLABDLDVKGQNGHMGHLHYSVRLVENVIMVLVKFPGVYKVLNLYCHSLIAQLQIAY 391
Db 298 LKIDNPFLISLQSONTVRLIYIYMMTRPENSVILLWYFKTDIYMMVYCAPLILQLIGY 357

Qy 392 LISIDMFLPQQYLHPLRSLETHNYVD---YLHCVYCHQHPRTRYVENSEP 438
Db 358 CTGILFMLVYIOPFHCKKLFSSSVSESFALLRACWSS--LRRKSSEP 405

RESULT 5

Sequence 33387, Application US/09864761
Patent No. US20020048761A1

GENERAL INFORMATION:

APPLICANT: Penn, Sharron G.

APPLICANT: Rank, David R.

APPLICANT: Hanzel, David K.

APPLICANT: Chen, Wenheng

TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR MICROARRAY

CURRENT APPLICATION NUMBER: US/09/864,761

CURRENT FILING DATE: 2001-05-23

PRIOR APPLICATION NUMBER: US 60/180,312

PRIOR FILING DATE: 2000-02-04

PRIOR APPLICATION NUMBER: US 60/207,456

PRIOR APPLICATION NUMBER: US 09/632,366

PRIOR FILING DATE: 2000-08-03

PRIOR APPLICATION NUMBER: GB 24263,6

PRIOR FILING DATE: 2000-10-04

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR FILING DATE: 2000-09-27

PRIOR APPLICATION NUMBER: PCT/US01/00664

PRIOR FILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00666

PRIOR FILING DATE: 2001-01-10

PRIOR APPLICATION NUMBER: PCT/US01/00668

PRIOR FILING DATE: 2001-01-10

PRIOR APPLICATION NUMBER: PCT/US01/00663

PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00662
PRIOR FILING DATE: 2000-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00687
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annotrax Sequence Listing Engine vers. 1.1
SEQ ID NO: 33387
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
OTHER INFORMATION: MAP TO AC005301.16
OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 1.3
OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 1.5
OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 1.5
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 1.3
OTHER INFORMATION: EXPRESSED IN HBL100, SIGNAL = 1.2
OTHER INFORMATION: EXPRESSED IN HELIX, SIGNAL = 1.7
OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 1.6
OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL = 1.1
OTHER INFORMATION: EXPRESSED IN RETAL LIVER, SIGNAL = 1.3
OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.8
OTHER INFORMATION: EST HUMAN HIT: AA256009_1, EVALUATE 3.00e-13
OTHER INFORMATION: SWISSPROT HIT: P51811, EVALUATE 8.00e-41
US-09-864-751-33387

Query Match 26.2%; Score 607; DB 10; Length 216;
Best Local Similarity 55.1%; Prod. No. 2.5e-50;
Matches 109; Conservative 42; Mismatches 47; Indels 0; Gaps 0;

Qy 213 TYGATCNMLAQIKYDDYKIRLGPLVLCITIWRTEBITSRLLVLFSATLRLKAVP 272
Db 1 TYGATCNMLAQISNDTTKLPPBFFCVMWRFLEVIRVTLAFFIASLKLKSLPV 60

Qy 273 LYLNFLITLFEFWKMRSGAMPNNTKBNFVGTLLVVLISVTLIYAGINFSCSWSALQ 332
Db 61 LLIJYFVSLLAWLFWKSGAHLPGNBENNSNNVGTVMLFLITLAAINFSCSWSAYL 120

Qy 333 RLADRLDVKGONWGHMGHLHYSVRLVENVIMVLVKFPGVYKVLNLYCHSLIAQLIAYL 392
Db 121 QLSDDKLIDGRQRGRHLHYSFQFLNCCDSLIAVOLISYL 180

Qy 393 ISIDFMFLFFYFLPHPLRS 410
Db 181 LATGMFLFYQLYPHQS 198

RESULT 6

US-10-092-154-595
Sequence 595, Application US/10092154
Publication No. US20030054375A1
GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PCT09C1
CURRENT APPLICATION NUMBER: US/10/092,154
CURRENT FILING DATE: 2002-03-07
NUMBER OF SEQ ID NOS: 2003
Prior Application removed - See File Wrapper or Palm
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO: 595
LENGTH: 125
TYPE: PRT
ORGANISM: Homo sapiens

FEATURE:
 NAME/KEY: misc_feature
 LOCATION: (70)
 OTHER INFORMATION: xaa equals any of the naturally occurring L-amino acids
 NAME/KEY: misc_feature
 LOCATION: (75)
 OTHER INFORMATION: xaa equals any of the naturally occurring L-amino acids
 NAME/KEY: misc_feature
 LOCATION: (81)
 OTHER INFORMATION: xaa equals any of the naturally occurring L-amino acids
 NAME/KEY: misc_feature
 LOCATION: (83)
 OTHER INFORMATION: xaa equals any of the naturally occurring L-amino acids
 US-10-092-154-595

Query Match 24.7%; Score 572; DB 9; Length 125;
 Best Local Similarity 94.7%; Pred. No. 2.6e-47;
 Matches 108; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

Qy 312 LISVTILYAGINPFCWSAQLRLADRIVDKGQNMGHLHYSLTFLNVDYLHCVCC 371
 Db 12 LISVTILYAGINPFCWSAQLRLADRIVDKGQNMGHLHYSLTFLNVDYLHCVCC 71

Qy 372 VVKVLYNCHSLIAQLIIAYLISDEMILFFQVLPRLSLFTHNVDYLHCVCC 425
 Db 72 VVKVLYNCHSLXKLAQLIIAYLISDEMILFFQVLPRLSLFTHNVDYLHCVCC 125

RESULT 7
 US-09-764-847-595
 Sequence 595, Application US/09764847
 GENERAL INFORMATION:
 APPLICANT: Roben et al.
 TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
 FILE REFERENCE: PC009
 CURRENT APPLICATION NUMBER: US/09/764, 847
 CURRENT FILING DATE: 2001-01-17
 PRIOR APPLICATION data removed - consult PALM or file wrapper
 NUMBER OF SEQ ID NOS: 2003
 SOFTWARE: PatentIn Ver. 2.0
 SEQ ID NO: 595
 LENGTH: 125
 TYPE: PRT
 ORGANISM: Homo sapiens
 FEATURE:
 NAME/KEY: SITE
 LOCATION: (70)
 OTHER INFORMATION: xaa equals any of the naturally occurring L-amino acids
 NAME/KEY: SITE
 LOCATION: (75)
 OTHER INFORMATION: xaa equals any of the naturally occurring L-amino acids
 NAME/KEY: SITE
 LOCATION: (81)
 OTHER INFORMATION: xaa equals any of the naturally occurring L-amino acids
 NAME/KEY: SITE
 LOCATION: (83)
 OTHER INFORMATION: xaa equals any of the naturally occurring L-amino acids
 US-09-764-847-595

Query Match 24.7%; Score 572; DB 10; Length 125;
 Best Local Similarity 94.7%; Pred. No. 2.6e-47;
 Matches 109; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

RESULT 8
 US-09-864-761-37894
 Sequence 37894, Application US/09864761
 Patent No. US200301048763A1
 GENERAL INFORMATION:
 APPLICANT: Penn, Sharron G.
 APPLICANT: Rank, David R.
 APPLICANT: Hanzel, David K.
 APPLICANT: Chen, Wansheng
 TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR MICROARRAY
 FILE REFERENCE: Acomica-X-1
 CURRENT APPLICATION NUMBER: US/09/864, 761
 CURRENT FILING DATE: 2001-05-23
 PRIOR APPLICATION NUMBER: US 60/180,312
 PRIOR FILING DATE: 2000-02-04
 PRIOR APPLICATION NUMBER: US 60/207,456
 PRIOR FILING DATE: 2000-05-26
 PRIOR APPLICATION NUMBER: US 09/632,366
 PRIOR FILING DATE: 2000-08-03
 PRIOR APPLICATION NUMBER: GB 24263, 6
 PRIOR FILING DATE: 2000-10-04
 PRIOR APPLICATION NUMBER: PCT/US01/00666
 PRIOR FILING DATE: 2000-09-27
 PRIOR APPLICATION NUMBER: PCT/US01/00666
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00667
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00664
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00669
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00665
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00668
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00663
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00662
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00661
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: PCT/US01/00670
 PRIOR FILING DATE: 2001-01-30
 PRIOR APPLICATION NUMBER: US 60/234, 687
 PRIOR FILING DATE: 2000-09-21
 PRIOR APPLICATION NUMBER: US 09/608, 408
 PRIOR FILING DATE: 2000-06-30
 PRIOR APPLICATION NUMBER: US 09/774, 203
 NUMBER OF SEQ ID NOS: 49117
 SOFTWARE: Anomax Sequence Listing Engine vers. 1.1
 SEQ ID NO: 37894
 TYPE: PRT
 ORGANISM: Homo sapiens
 FEATURE:
 OTHER INFORMATION: MAP TO AC007064.22
 Best Local Similarity 59.6%; Pred. No. 1.8e-26;
 Matches 65; Conservative 23; Mismatches 21; Indels 0; Gaps 0;

Query Match 15.3%; Score 354; DB 10; Length 128;
 OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 1.1
 OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 0.92
 OTHER INFORMATION: EST HUMAN HIT: A225609.1, EVALU 2.0e-13
 OTHER INFORMATION: SWISSPROT HIT: P51811, EVALU 5.00e-18
 US-09-864-761-37894

Qy 312 LISVTILYAGINPFCWSAQLRLADRIVDKGQNMGHLHYSLTFLNVDYLHCVCC 371
 Db 12 LISVTILYAGINPFCWSAQLRLADRIVDKGQNMGHLHYSLTFLNVDYLHCVCC 71

Qy 372 VVKVLYNCHSLIAQLIIAYLISDEMILFFQVLPRLSLFTHNVDYLHCVCC 425
 Db 72 VVKVLYNCHSLXKLAQLIIAYLISDEMILFFQVLPRLSLFTHNVDYLHCVCC 125

Qy 302 NFSRPGTGLVVLISVTILYAGINPFCWSAQLRLADRIVDKGQNMGHLHYSLTFLNVDYLHCVCC 361
 Db 2 NSNNVGTMLFLITLIAAINFSCWSAVKLOSDDKLIDGRWGHRLHYSPFLENV 61

Qy 362 IMVLVFKFVGKVLLNYCHSLIAQLIIAYLISDEMILFFQVLPRLSLFTHNVDYLHCVCC 410

Page 5

RESULT 11
 US-10-176-758-524
 Qy 388 II---AYLISIDPMILLFFQYFLHP 407
 Db 297 IAGVUSGFLIGSVSLLVIVYSSLLHP 320

RESULT 12
 US-10-175-737-524
 Qy 388 II---AYLISIDPMILLFFQYFLHP 407
 Db 297 IAGVUSGFLIGSVSLLVIVYSSLLHP 320

GENERAL INFORMATION:
 Sequence 524, Application US/10176758
 Publication No. US20030008353A1

APPLICANT: Baker, Kevin P.
 APPLICANT: Chen, Jian
 APPLICANT: Desnoyers, Luc
 APPLICANT: Goddard, Audrey
 APPLICANT: Godowski, Paul J.
 APPLICANT: Gurney, Austin L.
 APPLICANT: Pan, James
 APPLICANT: Smith, Victoria
 APPLICANT: Watanabe, Colin K.
 APPLICANT: Wood, William I.

TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC ACIDS ENCODING THE SAME

FILE REFERENCE: P3430RIC10
 CURRENT APPLICATION NUMBER: US/10/175,737
 CURRENT FILING DATE: 2002-06-19
 Prior Application removed - See File wrapper or Palm
 NUMBER OF SEQ ID NOS: 612

SEQ ID NO 524
 LENGTH: 686
 TYPE: PRT
 ORGANISM: Homo Sapien
 US-10-175-737-524

Query Match 4.9%; Score 113; DB 9; Length 686;
 Best Local Similarity 20.8%; Pred. No. 0.018;
 Matches 80; Conservative 57; Mismatches 159; Indels 88; Gaps 15;

Qy 40 STFLYCGEASALIYMRVYRKNSETYRMTYTFSSMFIYVQLTLFVHDLAKDKPLS 99
 Db 9 SALLQAAEQSARLYTVAAYFTGRILLWGLALAVLILPGLVQADSYLWFRADGPGHCSL 68

Qy 100 LFPHLILLGPIVRCLEAMIKYLTWKKKEQEPYVSLTRKRMILIDGEEVILLEVYGHISR 159
 Db 69 VMLHLQLQGWKRHWDAALTSL---QKELEAPHRG----W-----W----- 101

Qy 160 TLAMHNRAYKRMQIQARLQLGVPQLTOLYVLSAEPYPLGRVYMEVLSVTVYGAFLC 219
 Db 102 -LQLEADLSALRLLEALQTGPFLQLQTYVELASDFDIVPGYSTLFWSSLWAL--- 157

Qy 220 NMLAIQIKYDDYKIRLG-----LEVLCITIWRLTEITSRLILVLFSATLKLKAVPF 272
 Db 158 -----VSYTRFMFGFMKGFLAMPWAALFCQQQWRMGMGLGTRVSLVLF-----YKAYHF 206

Qy 273 LVLFNLFLFEPWI-KFWRSQAQMPPNNIEKNFS-RVGTLYVVLISVTLYAGINFSCWSA 329
 Db 207 WV---FVVAQAHWLWV-AQQSDIDSTCHWRMLFLVGAIVYILCP-----LSFWDS 258

Qy 330 LQLRLAARDLVQDQNHMGFLYSVRIVENVMVLUVKPF--GVKVLLNCHSIALQL 387
 Db 259 -----PSRNRMVTFYMMVMLLENILLLIATDFLQGA-----SWTSLSQT 296

Qy 388 II---AYLISIDPMILLFFQYFLHP 407
 Db 297 IAGVUSGFLIGSVSLLVIVYSSLLHP 320

RESULT 13
 US-10-173-706-524
 Qy 330 LQLRLAARDLVQDQNHMGFLYSVRIVENVMVLUVKPF--GVKVLLNCHSIALQL 387
 Db 259 -----PSRNRMVTFYMMVMLLENILLLIATDFLQGA-----SWTSLSQT 296

GENERAL INFORMATION:
 Application No. US2003002293A1
 Publication No. US2003002293A1
 Baker, Kevin P.
 Chen, Jian
 Desnoyers, Luc
 Goddard, Audrey
 Godowski, Paul J.
 Gurney, Austin L.
 Pan, James
 Smith, Victoria
 Watanabe, Colin K.
 Wood, William I.

TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC ACIDS ENCODING THE SAME

FILE REFERENCE: P3430RIC10
 CURRENT APPLICATION NUMBER: US/10173706
 CURRENT FILING DATE: 2002-06-19
 Prior Application removed - See File wrapper or Palm
 NUMBER OF SEQ ID NOS: 612

SEQ ID NO 524
 LENGTH: 686
 TYPE: PRT
 ORGANISM: Homo Sapien
 US-10-173-706-524

Query Match 4.9%; Score 113; DB 9; Length 686;
 Best Local Similarity 20.8%; Pred. No. 0.018;
 Matches 80; Conservative 57; Mismatches 159; Indels 88; Gaps 15;

Qy 40 STFLYCGEASALIYMRVYRKNSETYRMTYTFSSMFIYVQLTLFVHDLAKDKPLS 99
 Db 9 SALLQAAEQSARLYTVAAYFTGRILLWGLALAVLILPGLVQADSYLWFRADGPGHCSL 68

Qy 100 LFPHLILLGPIVRCLEAMIKYLTWKKKEQEPYVSLTRKRMILIDGEEVILLEVYGHISR 159
 Db 69 VMLHLQLQGWKRHWDAALTSL---QKELEAPHRG----W-----W----- 101

Qy 160 TLAMHNRAYKRMQIQARLQLGVPQLTOLYVLSAEPYPLGRVYMEVLSVTVYGAFLC 219
 Db 102 -LQLEADLSALRLLEALQTGPFLQLQTYVELASDFDIVPGYSTLFWSSLWAL--- 157

Qy 220 NMLAIQIKYDDYKIRLG-----LEVLCITIWRLTEITSRLILVLFSATLKLKAVPF 272
 Db 158 -----VSYTRFMFGFMKGFLAMPWAALFCQQQWRMGMGLGTRVSLVLF-----YKAYHF 206

Qy 273 LVLFNLFLFEPWI-KFWRSQAQMPPNNIEKNFS-RVGTLYVVLISVTLYAGINFSCWSA 329
 Db 207 WV---FVVAQAHWLWV-AQQSDIDSTCHWRMLFLVGAIVYILCP-----LSFWDS 258

Qy 330 LQLRLAARDLVQDQNHMGFLYSVRIVENVMVLUVKPF--GVKVLLNCHSIALQL 387
 Db 259 -----PSRNRMVTFYMMVMLLENILLLIATDFLQGA-----SWTSLSQT 296

Qy 388 II---AYLISIDPMILLFFQYFLHP 407
 Db 297 IAGVUSGFLIGSVSLLVIVYSSLLHP 320

RESULT 14
 US-10-173-706-524
 Qy 330 LQLRLAARDLVQDQNHMGFLYSVRIVENVMVLUVKPF--GVKVLLNCHSIALQL 387
 Db 259 -----PSRNRMVTFYMMVMLLENILLLIATDFLQGA-----SWTSLSQT 296

GENERAL INFORMATION:
 Application No. US2003002293A1
 Publication No. US2003002293A1
 Baker, Kevin P.
 Chen, Jian
 Desnoyers, Luc
 Goddard, Audrey
 Godowski, Paul J.
 Gurney, Austin L.
 Pan, James
 Smith, Victoria
 Watanabe, Colin K.
 Wood, William I.

TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC ACIDS ENCODING THE SAME

FILE REFERENCE: P3430RIC10
 CURRENT APPLICATION NUMBER: US/10173706
 CURRENT FILING DATE: 2002-06-19
 Prior Application removed - See File wrapper or Palm
 NUMBER OF SEQ ID NOS: 612

SEQ ID NO 524
 LENGTH: 686
 TYPE: PRT
 ORGANISM: Homo Sapien
 US-10-173-706-524

Query Match 4.9%; Score 113; DB 9; Length 686;
 Best Local Similarity 20.8%; Pred. No. 0.018;
 Matches 80; Conservative 57; Mismatches 159; Indels 88; Gaps 15;

Qy 40 STFLYCGEASALIYMRVYRKNSETYRMTYTFSSMFIYVQLTLFVHDLAKDKPLS 99
 Db 9 SALLQAAEQSARLYTVAAYFTGRILLWGLALAVLILPGLVQADSYLWFRADGPGHCSL 68

Qy 100 LFPHLILLGPIVRCLEAMIKYLTWKKKEQEPYVSLTRKRMILIDGEEVILLEVYGHISR 159
 Db 69 VMLHLQLQGWKRHWDAALTSL---QKELEAPHRG----W-----W----- 101

Qy 160 TLAMHNRAYKRMQIQARLQLGVPQLTOLYVLSAEPYPLGRVYMEVLSVTVYGAFLC 219
 Db 102 -LQLEADLSALRLLEALQTGPFLQLQTYVELASDFDIVPGYSTLFWSSLWAL--- 157

Qy 220 NMLAIQIKYDDYKIRLG-----LEVLCITIWRLTEITSRLILVLFSATLKLKAVPF 272
 Db 158 -----VSYTRFMFGFMKGFLAMPWAALFCQQQWRMGMGLGTRVSLVLF-----YKAYHF 206

Qy 273 LVLFNLFLFEPWI-KFWRSQAQMPPNNIEKNFS-RVGTLYVVLISVTLYAGINFSCWSA 329
 Db 207 WV---FVVAQAHWLWV-AQQSDIDSTCHWRMLFLVGAIVYILCP-----LSFWDS 258

Qy 330 LQLRLAARDLVQDQNHMGFLYSVRIVENVMVLUVKPF--GVKVLLNCHSIALQL 387
 Db 259 -----PSRNRMVTFYMMVMLLENILLLIATDFLQGA-----SWTSLSQT 296

Qy 388 II---AYLISIDPMILLFFQYFLHP 407
 Db 297 IAGVUSGFLIGSVSLLVIVYSSLLHP 320

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; TITLE OF INVENTION: ACIDS ENCODING THE SAME
; FILE REFERENCE: P3430R1C7
; CURRENT APPLICATION NUMBER: US/10/173, 706
; CURRENT FILING DATE: 2002-06-17
; Prior Application removed - See File Wrapper or Palm
; NUMBER OF SEQ ID NOS: 612
; SEQ ID NO 524
; LENGTH: 686
; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-173-706-524

Query Match Score 113; DB 9; Length 686
Best Local Similarity 20.8%; Pred. No. 0.018;
Matches 80; Conservative 57; Mismatches 159; Identity 524
; SEQ ID NO 524
; LENGTH: 686
; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-173-706-524

Qy 40 STFLYCEBAASALYMLTYRKNSETYRMITYTFSPEMFSSSIMVQLTL
Db 9 SALLQAEQSARLYTVAYFTGRLLGWLALAVLPGPVQAASY
Qy 100 LPMHLILLGPIRCLEAMIKYLTWKEQEOPPYVSLTRKKMFLDGG
Db 69 VMLHLLQLGWNRKHWDAALTSL---QKCLEAPHRG
Qy 160 TLAMHRNAYKRMQSQIQFLGSVQLTYOLYVLSLISAEVPLGRVYLM
Db 102 -LQLQEDALSARLLEALLQTGHLLQTYVFLASDFTD1IVPGVST
Qy 220 NMLAIQSKYDDYKIRLGP-----LEVLCITIWRTLETSRLJL
Db 158 -----VSYTRNGFMKGHLAMPWAALFCQQLWRMGMLCTRVLISL
Qy 273 LVLNPLILPFWI--KFWRSQGQMPPNNIEKNF5-RVGFLVVLISV
Db 207 WV---FVVAQAHWLVMTFWLV-AQSDSLIDSTCHWRFLNIVGAVY
Qy 330 LQLRLADRLVDKGQNNGMGLHYSVRLVENVIMVLVFKFEP--GVYK
Db 259 -----PSRNRMVTYPMVMLENILILLALATDFLQGA-
Qy 388 II----AYLISIDFMLFFQYLHP 407
Db 297 IAGVLSGFLIGSVSLLVYSSLHP 320

RESULT 14
US-10-175-738-524
; Sequence 524, Application US/10/175,738
; Publication No. US2003002294A1
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin P.
; APPLICANT: Chen, Jian
; APPLICANT: Desnoyers, Luc
; APPLICANT: Goddard, Audrey
; APPLICANT: Godebski, Paul J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Pan, James
; APPLICANT: Smith, Victoria
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDE
; FILE REFERENCE: P3430R1C45
; CURRENT APPLICATION NUMBER: US/10/175, 738
; CURRENT FILING DATE: 2002-06-19
; Prior Application removed - See File Wrapper or Palm
; NUMBER OF SEQ ID NOS: 612
; SEQ ID NO 524
; LENGTH: 686
; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-175-738-524

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Query Match          4.0%; Score 113; DB 9; Length 686;
Best Local Similarity 20.8%; Pred. No. 0.018;
Matches 80; Conservative 57; Mismatches 159; Indels 88; Gaps 15;
  40 STEFLYGEAAASALYMRVIRKNSETRYMTYTFSEPMESSIMVQTLIFVHRLDAKDPLS 99
  9 SALLQAEQARLTVAYYFTGRLLGWGNALAVLPGFIVQALSYLWPAFDRGPHGHCSL 68
  100 LFMHLILLGIVRCLEAMIKYLTLWKKEDEEPYVSLTRKMLIDGEEVILEWEVGHISR 159
  69 VMLHLILQLGTWKRHDAALTSL---QKCELEAHPG-----W-----W----- 101
  160 TLAMHRNAYRGRMSQIQAFLGSVPQTYQLVSLISAEVPPGRVVLIVFSIVSVTYGATLIC 219
  102 -LQLQEAIDLSSALRLLAEALLQTGPILLQTVFLASDFTD1VPGYSTLFSWSSLWAL-- 157
  220 NMLA10IQKYDDYKIRLGP-----LEVLTITMRTLEITSRLJLJLVLFEGATLK1KAVPFF 272
  158 -----VSYTRFMGMKPGMPGHMLAMPMAAFLFQOLWRMGMLGTRVLVLF-----YKAYHF 206
  273 LYLNFLLIELEPWI--KFWSGAQMPNNIENKNF- RVGTLVVLISVTILYAGINFSCWSA 329
  207 WV--FVVAGAHWWMTFNLV-AQOSDIIISTCHWLFLNLYGAVYILCY----LSFWDS 258
  330 LQLRLADRLDVDKGQNWGHWGLHYSVRLYENVNIMVLFKEF--GVKVLLNYCHSLLJALQI 387
  259 -----PSRNRMNTFTVMNLLENILLLATLFLQGA-----SWTSLQR 296
  388 II---AYLSSIDFMILPFQYLYHP 407
  297 TAGVLSGFLIGSVSLVIVYSLHHP 320

RESULT 15
US-10-175-752-524
; Sequence 524, Application US/101757524
; Publication No. US20030022295A1
; GENERAL INFORMATION:
;   APPLICANT: Baker, Kevin P.
;   APPLICANT: Chen, Jian
;   APPLICANT: Desnoyers, Luc
;   APPLICANT: Goddard, Audrey
;   APPLICANT: Godowski, Paul J.
;   APPLICANT: Gurney, Austin L.
;   APPLICANT: Pan, James
;   APPLICANT: Smith, Victoria
;   APPLICANT: Watanabe, Colin K.
;   APPLICANT: Wood, William I.
;   APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
; TITLE OF INVENTION: ACIDS ENCODING THE SAME
; FILE REFERENCE: B343081C60
; CURRENT APPLICATION NUMBER: US/10/175,752
; CURRENT FILING DATE: 2002-06-19
; Prior Application removed - See File Wrapper or Palm
; NUMBER OF SEQ ID NOS: 612
; SEQ ID NO 524
; LENGTH: 686
; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-175-752-524

Query Match          4.0%; Score 113; DB 9; Length 686;
Best Local Similarity 20.8%; Pred. No. 0.018;
Matches 80; Conservative 57; Mismatches 159; Indels 88; Gaps 15;
  40 STEFLYGEAAASALYMRVIRKNSETRYMTYTFSEPMESSIMVQTLIFVHRLDAKDPLS 99
  9 SALLQAEQARLTVAYYFTGRLLGWGNALAVLPGFIVQALSYLWPAFDRGPHGHCSL 68
  100 LFMHLILLGIVRCLEAMIKYLTLWKKEDEEPYVSLTRKMLIDGEEVILEWEVGHISR 159

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| | | | | | |
|----|-----|--|-----------|--------|-----|
| db | 69 | VNLHLQLQGwVKRHWDAALTSL-- | - - - - - | W----- | 101 |
| Dy | 160 | TLMHRYAKRMSOIQAFGLGSVPQTYVLSLISAEVPLGRVVLMLVFSVTVYGTATLC | 219 | -- | |
| db | 102 | : : : : : : : : : : : : : : : : | | | |
| Dy | 102 | -LQLEADLSALRLEALQTGPFLILLQTYVFLASDFTD1VPGSTLFSMSSLSVAL-- | 157 | | |
| db | 220 | NMLIAQIKYDDKYLKRLGP-----LEVLCITIWTITLEITSRLLLILVLPSATLKAVPF | 272 | -- | |
| Dy | 158 | : : : : : : : : : : : : : : : : | | | |
| db | 158 | --VSYTRENGFMKPGFLAMPWAALPQQLMRGMGLGTRVLSVLP----YKAYHF | 206 | | |
| Dy | 273 | LVLNLF1LFEPEI--KFWRSQAOPNNIEKE NFS--RGVTLLVLVLSITVLYAGINSFCWSA | 329 | -- | |
| db | 207 | : : : : : : : : : : : : : : : : | | | |
| Dy | 207 | WV--FVVAGAHPLVMTFLW-AQOSDIIDSTCHWLRFNLVGAIVLCY--LSPFDWS | 258 | | |
| db | 330 | LQLRLADRDLVDKGCQNQNGHMGHLHSVRVLYENVNIMVLPKF--GYKVLLNYCHSIALQL | 387 | -- | |
| Dy | 259 | : : : : : : : : : : : : : : : : | | | |
| db | 388 | --PSRNENVTPYVMLENILILLILLTDFLQGA-----SWTSLSQRT | 296 | | |
| Dy | 388 | II--AYLISIDFMMIIFPOYIHP | 407 | -- | |
| db | 297 | IAGVLSGPGISGVSYVLSLILH | 320 | | |

Search completed: April 1, 2003, 08:51:03
Job time : 21 SECS